SEQUENCE LISTING

- (1) GENERAL INFORMATION
- (ii) TITLE OF THE INVENTION: CTLA4/CD28Ig HYBRID FUSION PROTEINS AND USES THEREOF
- (iii) NUMBER OF SEQUENCES: 33
- (iv) CORRESPONDENCE ADDRESS:
 - (A) ADDRESSEE: Mandel & Adriano
 - (B) STREET: 35 N. Arroyo Parkway, Suite 60
 - (C) CITY: Pasadena
 - (D) STATE: California
 - (E) COUNTRY: USA
 - (F) ZIP: 91103
- (v) COMPUTER READABLE FORM:
- (A) MEDIUM TYPE: Diskette
- (B) COMPUTER: IBM Compatible
- (C) OPERATING SYSTEM: DOS
- (D) SOFTWARE: FastSEQ Version 2.0
- (vi) CURRENT APPLICATION DATA:
 - (A) APPLICATION NUMBER: 09/454,651
 - (B) FILING DATE: 06-DEC-1999
 - (C) CLASSIFICATION:
- (vii) PRIOR APPLICATION DATA:
 - (A) APPLICATION NUMBER: 08/228,208
 - (B) FILING DATE: 15-APR-1994
 - (A) APPLICATION NUMBER: 08/008,898
 - (B) FILING DATE: 22-JAN-1993
 - (A) APPLICATION NUMBER: 07/723,617
 - (B) FILING DATE: 27-JUL-1991
- (viii) ATTORNEY/AGENT INFORMATION:
 - (A) NAME: Adriano, Sarah B
 - (B) REGISTRATION NUMBER: 34,470
 - (C) REFERENCE/DOCKET NUMBER: 30436.30USD1
 - (ix) TELECOMMUNICATION INFORMATION:
 - (A) TELEPHONE: 626 395-7801
 - (B) TELEFAX: 626 395-0694
 - (C) TELEX:



(2) INFORMATION FOR SEQ ID NO:1: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 39 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:	
CTAGCCACTG AAGCTTCACC ATGGGTGTAC TGCTCACAC	39
(2) INFORMATION FOR SEQ ID NO:2: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 39 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:	
TGGCATGGGC TCCTGATCAG GCTTAGAAGG TCCGGGAAA	39
(2) INFORMATION FOR SEQ ID NO:3: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 39 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3: TTTGGGCTCC TGATCAGGAA AATGCTCTTG CTTGGTTGT	39
(2) INFORMATION FOR SEQ ID NO:4: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 84 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:	
AAGCAAGAGC ATTTTCCTGA TCAGGAGCCC AAATCTTCTG ACAAAACTCA CACATCCCCA	60
CCGTCCCCAG CACCTGAACT CCTG	84
(2) INFORMATION FOR SEQ ID NO:5: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 41 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	

	(ii) MOLECULE TYPE: DNA (genomic) (xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:	
CTTC	GACCAG TCTAGAAGCA TCCTCGTGCG ACCGCGAGAG C	41
	INFORMATION FOR SEQ ID NO:6: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 47 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) (xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:	47
CAT'I	GCACAG TCAAGCTTCC ATGCCCATGG GTTCTCTGGC CACCTTG	47
(2)	<pre>INFORMATION FOR SEQ ID NO:7: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 39 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) (xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:</pre>	
ATCO	CACAGTG CAGTGATCAT TTGGATCCTG GCATGTGAC	39
(2)	<pre>INFORMATION FOR SEQ ID NO:8: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 65 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) (xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:</pre>	
CTC	AGTCTGG TCCTTGCACT CCTGTTTCCA AGCATGGCGA GCATGGCAAT GCACGTGGCC	60
CAG	cc	65
(2)	<pre>INFORMATION FOR SEQ ID NO:9: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 33 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) (xi) SEQUENCE DESCRIPTION: SEQ ID NO:9:</pre>	
TTT	GGGCTCC TGATCAGAAT CTGGGCACGG TTG	33

(2) INFORMATION FOR SEQ ID NO:10:

(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 72 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) (xi) SEQUENCE DESCRIPTION: SEQ ID NO:10: CTAGCCACTG AAGCTTCACC AATGGGTGTA CTGCTCACAC AGAGGACGCT GCTCAGTCTG	60											
GTCCTTGCAC TC												
(2) INFORMATION FOR SEQ ID NO:11: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 33 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) (xi) SEQUENCE DESCRIPTION: SEQ ID NO:11: GCAATGCACG TGGCCCAGCC TGCTGTGGTA GTG	33											
(2) INFORMATION FOR SEQ ID NO:12: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 45 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) (xi) SEQUENCE DESCRIPTION: SEQ ID NO:12: TGATGTAACA TGTCTAGATC AATTGATGGG AATAAAATAA	45											
(2) INFORMATION FOR SEQ ID NO:13: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 561 nucleotides (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) (ix) FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 1561												
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:13: GCA ATG CAC GTG GCC CAG CCT GCT GTG GTA CTG GCC AGC AGC CGA GGC GCA ATG CAC GTG GCC CAG CCT GCT GTG GTA CTG GCC AGC AGC CGA GGC	48											
GCA ATG CAC GTG GCC CAG CCT GCT GTG GTA CTG GGG STA CTG GTG GTA CTG GTG GTA CTG GTA CT												
ATC GCC AGC TTT GTG TGT GAG TAT GCA TCT CCA GGC AAA GCC ACT GAG Ile Ala Ser Phe Val Cys Glu Tyr Ala Ser Pro Gly Lys Ala Thr Glu 20 25 30	96											

'DB

GT(Va]	C CGG Arg	GTO Val	. Thr	GTG Val	CTT Leu	CGG Arg	CAG Gln 40	Ala	GAC Asp	AGC Ser	CAG Gln	GTG Val	Thr	GAA Glu	GTC Val	144
TG1 Cys	GCG Ala 50	АТа	ACC Thr	TAC	ATG Met	ATG Met 55	GGG Gly	AAT Asn	' GAG Glu	TTG Leu	ACC Thr 60	TTC Phe	CTA Leu	GAT Asp	GAT Asp	192
TCC Ser 65	тте	TGC Cys	ACG Thr	GGC Gly	ACC Thr 70	Ser	AGT Ser	GGA Gly	AAT Asn	CAA Gln 75	GTG Val	AAC Asn	CTC Leu	ACT Thr	ATC Ile 80	240
CAA Gln	GGA Gly	CTG Leu	AGG Arg	GCC Ala 85	ATG Met	GAC Asp	ACG Thr	GGA Gly	CTC Leu 90	TAC Tyr	ATC Ile	TGC Cys	AAG Lys	GTG Val 95	GAG Glu	288
CTC Leu	ATG Met	TAC Tyr	CCA Pro 100	CCG Pro	CCA Pro	TAC Tyr	TAC Tyr	CTG Leu 105	GGC Gly	ATA Ile	GGC Gly	AAC Asn	GGA Gly 110	ACC Thr	CAG Gln	336
ATT Ile	TAT Tyr	GTA Val 115	ATT Ile	GAT Asp	CCA Pro	GAA Glu	CCG Pro 120	TGC Cys	CCA Pro	GAT Asp	TCT Ser	GAC Asp 125	TTC Phe	CTC Leu	CTC Leu	384
TGG Trp	ATC Ile 130	CTT Leu	GCA Ala	GCA Ala	GTT Val	AGT Ser 135	TCG Ser	GGG Gly	TTG Leu	TTT Phe	TTT Phe 140	TAT Tyr	AGC Ser	TTT Phe	CTC Leu	432
CTC Leu 145	ACA Thr	GCT Ala	GTT Val	TCT Ser	TTG Leu 150	AGC Ser	AAA Lys	ATG Met	CTA Leu	AAG Lys 155	AAA Lys	AGA Arg	AGC Ser	CCT Pro	CTT Leu 160	480
ACA Thr	ACA Thr	GGG Gly	GTC Val	TAT Tyr 165	GTG Val	AAA Lys	ATG Met	CCC Pro	CCA Pro 170	ACA Thr	GAG Glu	CCA Pro	Glu	TGT Cys 175	GAA Glu	528
AAG Lys	CAA Gln	TTT Phe	CAG Gln 180	CCT Pro	TAT Tyr	TTT . Phe	Ile	CCC Pro 185	ATC Ile	AAT Asn						561

(2) INFORMATION FOR SEQ ID NO:14:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 187 amino acids
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: protein
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:14:

Ala Met His Val Ala Gln Pro Ala Val Val Leu Ala Ser Ser Arg Gly

1 15

Ile Ala Ser Phe Val Cys Glu Tyr Ala Ser Pro Gly Lys Ala Thr Glu
20 25 30

Val Arg Val Thr Val Leu Arg Gln Ala Asp Ser Gln Val Thr Glu Val



	40	4
35	10	

	_ •		mla se	Tyr	Met	Met	Glv	Asn	Glu	Leu	Thr	Phe	Leu	Asp	Asp
Cys	Ala	Ala	Thr	тÀт	Mec		021				60				
	50					55					•				

- Ser Ile Cys Thr Gly Thr Ser Ser Gly Asn Gln Val Asn Leu Thr Ile 65 70 75 80
- Gln Gly Leu Arg Ala Met Asp Thr Gly Leu Tyr Ile Cys Lys Val Glu 85 90 95
- Leu Met Tyr Pro Pro Pro Tyr Tyr Leu Gly Ile Gly Asn Gly Thr Gln
 100 105 110
- Ile Tyr Val Ile Asp Pro Glu Pro Cys Pro Asp Ser Asp Phe Leu Leu 115 120 125
- Trp Ile Leu Ala Ala Val Ser Ser Gly Leu Phe Phe Tyr Ser Phe Leu 130 135 140
- Leu Thr Ala Val Ser Leu Ser Lys Met Leu Lys Lys Arg Ser Pro Leu 145 150 155 160
- Thr Thr Gly Val Tyr Val Lys Met Pro Pro Thr Glu Pro Glu Cys Glu 165 170 175

Lys Gln Phe Gln Pro Tyr Phe Ile Pro Ile Asn 180 185

(2) INFORMATION FOR SEQ ID NO:15:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 18 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: DNA (genomic)
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:15:

AATACGACTC ACTATAGG

18

(2) INFORMATION FOR SEQ ID NO:16:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 18 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: DNA (genomic)
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:16:

CACCACACTG TATTAACC

18

(2) INFORMATION FOR SEQ ID NO:17:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 223 amino acids

- (B) TYPE: amino acid
- (C) STRANDEDNESS: unknown
- (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: protein
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:17:

Met Ala Cys Leu Gly Phe Gln Arg His Lys Ala Gln Leu Asn Leu Ala 1 5 10 15

Ala Arg Thr Trp Pro Cys Thr Leu Leu Phe Phe Leu Leu Phe Ile Pro 20 25 30

Val Phe Cys Lys Ala Met His Val Ala Gln Pro Ala Val Val Leu Ala 35 40 45

Ser Ser Arg Gly Ile Ala Ser Phe Val Cys Glu Tyr Ala Ser Pro Gly 50 55 60

Lys Ala Thr Glu Val Arg Val Thr Val Leu Arg Gln Ala Asp Ser Gln 65 70 75 80

Val Thr Glu Val Cys Ala Ala Thr Tyr Met Met Gly Asn Glu Leu Thr
85 90 95

Phe Leu Asp Asp Ser Ile Cys Thr Gly Thr Ser Ser Gly Asn Gln Val

Asn Leu Thr Ile Gln Gly Leu Arg Ala Met Asp Thr Gly Leu Tyr Ile 115 120 125

Cys Lys Val Glu Leu Met Tyr Pro Pro Pro Tyr Tyr Leu Gly Ile Gly 130 135 140

Asn Gly Thr Gln Ile Tyr Val Ile Asp Pro Glu Pro Cys Pro Asp Ser 145 150 155 160

Asp Phe Leu Leu Trp Ile Leu Ala Ala Val Ser Ser Gly Leu Phe Phe
165 170 175

Tyr Ser Phe Leu Leu Thr Ala Val Ser Leu Ser Lys Met Leu Lys Lys 180 185 190

Arg Ser Pro Leu Thr Thr Gly Val Tyr Val Lys Met Pro Pro Thr Glu
195 200 205

Pro Glu Cys Glu Lys Gln Phe Gln Pro Tyr Phe Ile Pro Ile Asn 210 215 220

- (2) INFORMATION FOR SEQ ID NO:18:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 223 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: unknown
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: protein
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:18:



Met 1	Ala	Cys	Leu	Gly 5	Leu	Arg	Arg	Tyr	Lys 10	Ala	Gln	Leu	Gln	Leu 15	Pro
Ser	Arg	Thr	Trp 20	Pro	Phe	Val	Ala	Leu 25	Leu	Thr	Leu	Leu	Phe 30	Ile	Pro
Val	Phe	Ser 35	Glu	Ala	Ile	Gln	Val 40	Thr	Gln	Pro	Ser	Val 45	Tyr	Leu	Ala
Ser	Ser 50	His	Gly	Tyr	Ala	Ser 55	Phe	Pro	Cys	Glu	Tyr 60	Ser	Pro	Ser	His
Asn 65	Thr	Asp	Glu	Val	Arg 70	Val	Thr	Val	Leu	Arg 75	Gln	Thr	Asn	Asp	Gln 80
Met	Thr	Glu	Val	Cys 85	Ala	Thr	Thr	Phe	Thr 90	Glu	Lys	Asn	Thr	Val 95	Gly
Phe	Leu	Asp	Tyr 100	Pro	Phe	Cys	Ser	Gly 105	Thr	Phe	Asn	Glu	Ser 110	Arg	Val
Asn	Leu	Thr 115	Ile	Gln	Gly	Leu	Arg 120	Ala	Val	Asp	Thr	Gly 125	Leu	Tyr	Leu
Cys	Lys 130	Val	Glu	Leu	Met	Tyr 135	Pro	Pro	Pro	Tyr	Phe 140	Val	Gly	Met	Gly
Asn 14		Thr	Gln	Ile	Tyr 15		Ile	Asp	Pro	Glu 15	Pro 5	Cys	Pro	Asp	Ser 160
Asp	Phe	Leu	Leu	Trp 165	Ile	Leu	Tyr	Ala	Val 170		Leu	Gly	Leu	Phe 175	Phe
Tyr	Ser	Phe	Leu 180		Ser	Ala	Val	Ser 185		Ser	Lys	Met	Leu 190	Lys	Lys
Arg	Ser	Pro		Thr	Thr	Gly	Val		Val	Lys	Met	Pro 205	Pro	Thr	Glu

(2) INFORMATION FOR SEQ ID NO:19:

210

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 218 amino acids

Pro Glu Cys Glu Lys Gln Phe Gln Pro Tyr Phe Ile Pro Ile Asn

- (B) TYPE: amino acid
- (C) STRANDEDNESS: unknown

215

- (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: protein
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:19:

Met Thr Leu Arg Leu Leu Phe Leu Ala Leu Asn Phe Phe Ser Val Gln
1 5 10 15

Val Thr Glu Asn Lys Ile Leu Val Lys Gln Ser Pro Leu Leu Tyr Val



20

25

30

Asp Ser Asn Glu Val Ser Leu Ser Cys Arg Tyr Ser Tyr Asn Leu Leu 35 40 45

Ala Lys Glu Phe Arg Ala Ser Leu Tyr Lys Gly Val Asn Ser Asp Val
50 55 60

Glu Val Cys Val Gly Asn Gly Asn Phe Thr Tyr Gln Pro Gln Phe Arg
65 70 75 80

Ser Asn Ala Glu Phe Asn Cys Asp Gly Asp Phe Asp Asn Glu Thr Val 85 90 95

Thr Phe Arg Leu Trp Asn Leu His Val Asn His Thr Asp Ile Tyr Phe

Cys Lys Ile Glu Phe Met Tyr Pro Pro Pro Tyr Leu Asp Asn Glu Arg

Ser Asn Gly Thr Ile Ile His Ile Lys Glu Lys His Leu Cys His Thr 130 135 140

Leu Phe Cys Tyr Gly Leu Leu Val Thr Val Ala Leu Cys Val Ile Trp 165 170 175

Thr Asn Ser Arg Arg Asn Arg Leu Leu Gln Val Thr Tyr Met Asn Met 180 185 190

Thr Pro Arg Arg Pro Gly Leu Thr Arg Lys Pro Tyr Gln Pro Tyr Ala 195 200 205

Pro Ala Arg Asp Phe Ala Ala Tyr Arg Pro 210 215

(2) INFORMATION FOR SEQ ID NO:20:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 218 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: unknown
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: protein
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:20:

Met Thr Leu Arg Leu Leu Phe Leu Ala Leu Ser Phe Phe Ser Val Gln
1 5 10 15

Val Thr Glu Asn Lys Ile Leu Val Lys Gln Ser Pro Leu Leu Val Tyr 20 25 30

Asp Asn Asn Glu Val Ser Leu Ser Cys Arg Tyr Ser Tyr Asn Leu Leu

Ala Lys Glu Phe Arg Ala Ser Leu Tyr Lys Gly Val Asn Ser Asp Val



50 55

Glu Val Cys Val Gly Asn Gly Asn Phe Thr Tyr Gln Pro Gln Phe Arg
65 70 75 80

60

Pro Asn Val Gly Phe Asn Cys Asp Gly Asn Phe Asp Asn Glu Thr Val 85 90 95

Thr Phe Arg Leu Trp Asn Leu Asp Val Asn His Thr Asp Ile Tyr Phe
100 105 110

Cys Lys Ile Glu Val Met Tyr Pro Pro Pro Tyr Leu Asp Asn Glu Lys 115 120 125

Ser Asn Gly Thr Ile Ile His Ile Lys Glu Lys His Leu Cys His Ala 130 135 140

Gln Thr Ser Pro Lys Leu Phe Trp Pro Leu Val Val Val Ala Gly Val
145 150 155 160

Leu Leu Cys Tyr Gly Leu Leu Tyr Thr Val Thr Leu Cys Ile Ile Trp
165 170 175

Thr Asn Ser Arg Arg Asn Arg Leu Leu Gln Ser Asp Tyr Met Asn Met 180 185 190

Thr Pro Arg Arg Leu Gly Pro Thr Arg Lys His Tyr Gln Pro Tyr Ala 195 200 205

Pro Ala Arg Asp Phe Ala Ala Tyr Arg Pro 210 215

(2) INFORMATION FOR SEQ ID NO:21:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 220 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: unknown
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: protein
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:21:

Met Leu Arg Leu Leu Leu Ala Leu Asn Leu Phe Pro Ser Ile Gln Val

Thr Gly Asn Lys Ile Leu Val Lys Gln Ser Pro Met Leu Val Ala Tyr 20 25 30

Asp Asn Ala Tyr Asn Leu Ser Cys Lys Tyr Ser Tyr Asn Leu Phe Ser 35 40 45

Arg Glu Phe Arg Ala Ser Leu His Lys Gly Leu Asp Ser Ala Val Glu 50 60

Val Cys Val Val Tyr Gly Asn Tyr Ser Gln Gln Leu Gln Val Tyr Ser 65 70 75 80



Lys Thr Gly Phe Asn Cys Asp Gly Lys Leu Gly Asn Glu Ser Val Thr 85 90 95

Phe Tyr Leu Gln Asn Leu Tyr Val Asn Gln Thr Asp Ile Tyr Phe Cys 100 105 110

Lys Ile Glu Val Met Tyr Pro Pro Pro Tyr Leu Asp Asn Glu Lys Ser 115 120 125

Asn Gly Thr Ile Ile His Val Lys Gly Lys His Leu Cys Pro Ser Pro 130 135 140

Gly Val Leu Ala Cys Tyr Ser Leu Leu Tyr Thr Val Ala Phe Ile Ile 165 170 175

Phe Trp Val Arg Ser Lys Arg Ser Arg Leu Leu His Ser Asp Tyr Met 180 185 190

Asn Met Thr Pro Arg Arg Pro Gly Pro Thr Arg Lys His Tyr Gln Pro 195 200 205

Tyr Ala Pro Pro Arg Asp Phe Ala Ala Tyr Arg Ser 210 215 220

(2) INFORMATION FOR SEQ ID NO:22:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 221 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: unknown
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: protein
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:22:

Met Leu Gly Ile Leu Val Val Leu Cys Leu Ile Pro Ala Ala Asp Val 1 5 10 15

Thr Glu Asn Lys Ile Leu Val Ala Gln Arg Pro Leu Leu Ile Val Ala 20 25 30

Asn Arg Thr Ala Thr Leu Val Cys Asn Tyr Thr Tyr Asn Gly Thr Gly 35 40 45

Lys Glu Phe Arg Ala Ser Leu His Lys Gly Thr Asp Ser Ala Val Glu 50 55 60

Val Cys Phe Ile Ser Trp Asn Met Thr Lys Ile Asn Ser Asn Ser Asn 65 70 75 80

Lys Glu Phe Asn Cys Arg Gly Ile His Asp Lys Asp Lys Val Ile Phe 85 90 95

Asn Leu Trp Asn Met Ser Ala Ser Gln Thr Asp Ile Tyr Phe Cys Lys
100 105 110

Ile Glu Ala Met Tyr Pro Pro Pro Tyr Val Tyr Asn Glu Lys Ser Asn 115 120 125

Gly Thr Val Ile His Tyr Arg Glu Thr Pro Ile Gln Thr Gln Glu Pro 130 135 140

Glu Ser Ala Thr Ser Tyr Trp Val Met Tyr Ala Val Thr Gly Leu Leu 145 150 155 160

Gly Phe Tyr Ser Met Leu Ile Thr Ala Val Phe Ile Ile Tyr Arg Gln 165 170 175

Lys Ser Lys Arg Asn Arg Tyr Arg Gln Ser Asp Tyr Met Asn Met Thr 180 185 190

Pro Arg His Pro Pro His Gln Lys Asn Lys Gly Tyr Pro Ser Tyr Ala 195 200 205

Pro Thr Arg Asp Tyr Thr Ala Tyr Arg Ser Trp Gln Pro 210 215 220

- (2) INFORMATION FOR SEQ ID NO:23:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 216 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: unknown
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: protein
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:23:

Gly Leu Ser His Phe Cys Ser Gly Val Ile His Val Thr Lys Glu Val 1 5 10

Lys Glu Val Ala Thr Leu Ser Cys Gly His Asn Val Ser Val Glu Glu 25 30

Leu Ala Gln Thr Arg Ile Tyr Trp Gln Lys Glu Lys Lys Met Val Leu 35 40 45

Thr Met Met Ser Gly Asp Met Asn Ile Trp Pro Glu Tyr Lys Asn Arg
50 55 60

Thr Ile Phe Asp Ile Thr Asn Asn Leu Ser Ile Val Ile Leu Ala Leu 65 70 75 80

Arg Pro Ser Asp Glu Gly Thr Tyr Glu Cys Val Val Leu Lys Tyr Glu
90
95

Lys Asp Ala Phe Lys Arg Glu His Leu Ala Glu Val Thr Leu Ser Val 100 105 110

Lys Ala Asp Phe Pro Thr Pro Ser Ile Ser Asp Phe Glu Ile Pro Thr 115 120 125

Ser Asn Ile Arg Arg Ile Ile Cys Ser Thr Ser Gly Gly Phe Pro Glu



140 135 130

Pro His Leu Ser Trp Leu Glu Asn Gly Glu Glu Leu Asn Ala Ile Asn 155 150

Thr Thr Val Ser Gln Asp Pro Glu Thr Glu Leu Tyr Ala Val Ser Ser 170

Lys Leu Asp Phe Asn Met Thr Thr Asn His Ser Phe Met Cys Leu Ile 185

Lys Tyr Gly His Leu Arg Val Asn Gln Thr Phe Asn Trp Asn Thr Thr 200 195

Lys Gln Glu His Phe Pro Asp Asn 210

- (2) INFORMATION FOR SEQ ID NO:24:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 6 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: unknown
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: protein
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:24:

Met Tyr Pro Pro Pro Tyr 1

(2) INFORMATION FOR SEQ ID NO:25:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 6 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: unknown
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: protein
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:25:

Ala Tyr Pro Pro Pro Tyr 1

- (2) INFORMATION FOR SEQ ID NO:26:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 6 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: unknown
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: protein
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:26:

Met Ala Pro Pro Pro Tyr 5

(A) LENGTH: 6 amino acids (B) TYPE: amino acid (C) STRANDEDNESS: unknown (D) TOPOLOGY: linear (ii) MOLECULE TYPE: protein (xi) SEQUENCE DESCRIPTION: SEQ ID NO:27: Met Tyr Ala Pro Pro Tyr (2) INFORMATION FOR SEQ ID NO:28: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 6 amino acids (B) TYPE: amino acid (C) STRANDEDNESS: unknown (D) TOPOLOGY: linear (ii) MOLECULE TYPE: protein (xi) SEQUENCE DESCRIPTION: SEQ ID NO:28: Met Tyr Pro Ala Pro Tyr (2) INFORMATION FOR SEQ ID NO:29: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 6 amino acids (B) TYPE: amino acid (C) STRANDEDNESS: unknown (D) TOPOLOGY: linear (ii) MOLECULE TYPE: protein (xi) SEQUENCE DESCRIPTION: SEQ ID NO:29: Met Tyr Pro Pro Ala Tyr (2) INFORMATION FOR SEQ ID NO:30: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 6 amino acids (B) TYPE: amino acid (C) STRANDEDNESS: unknown (D) TOPOLOGY: linear (ii) MOLECULE TYPE: protein (xi) SEQUENCE DESCRIPTION: SEQ ID NO:30: Met Tyr Pro Pro Pro Ala 5 1 (2) INFORMATION FOR SEQ ID NO:31:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 6 amino acids(B) TYPE: amino acid(C) STRANDEDNESS: unknown

(2) INFORMATION FOR SEQ ID NO:27:

(i) SEQUENCE CHARACTERISTICS:

```
Ala Ala Pro Pro Pro Tyr
  1
                  5
 (2) INFORMATION FOR SEQ ID NO:32:
       (i) SEQUENCE CHARACTERISTICS:
             (A) LENGTH: 6 amino acids
             (B) TYPE: amino acid
             (C) STRANDEDNESS: unknown
             (D) TOPOLOGY: linear
      (ii) MOLECULE TYPE: protein
      (xi) SEQUENCE DESCRIPTION: SEQ ID NO:32:
Met Tyr Pro Pro Ala Tyr
 1
                 5
(2) INFORMATION FOR SEQ ID NO:33:
       (i) SEQUENCE CHARACTERISTICS:
             (A) LENGTH: 6 amino acids
             (B) TYPE: amino acid
             (C) STRANDEDNESS: unknown
             (D) TOPOLOGY: linear
      (ii) MOLECULE TYPE: protein
      (xi) SEQUENCE DESCRIPTION: SEQ ID NO:33:
Met Tyr Pro Pro Pro Ala
  1 5
```

(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:31:

(ii) MOLECULE TYPE: protein